RAW SEQUENCE LISTING

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Application Serial Number: 10/561,906
Source: /FWD
Date Processed by STIC: 3/28/07

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RAW SEQUENCE LISTING   DATE: 03/28/2007
PATENT APPLICATION: US/10/561,906   TIME: 14:52:59.

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 3 <110> APPLICANT: Patek, Miroslav
 4         Elisakova, Veronika
 6 <120> TITLE OF INVENTION: FEEDBACK RESISTANT ACETOHYDROXY ACID SYNTHETHASE MUTANTS
 8 <130> FILE REFERENCE: 281642US0XPCT
10 <140> CURRENT APPLICATION NUMBER: US 10/561,906
11 <141> CURRENT FILING DATE: 2005-12-21
13 <150> PRIOR APPLICATION NUMBER: PCT/EP04/06157
14 <151> PRIOR FILING DATE: 2004-06-08
16 <150> PRIOR APPLICATION NUMBER: EP 03014640.1
17 <151> PRIOR FILING DATE: 2003-06-26
19 <160> NUMBER OF SEQ ID NOS: 12
21 <170> SOFTWARE: PatentIn version 3.3
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 519
25 <212> TYPE: DNA
26 <213> ORGANISM: Artificial Sequence
28 <220> FEATURE:
29 <223> OTHER INFORMATION: Synthetic DNA
32 <220> FEATURE:
33 <221> NAME/KEY: CDS
34 <222> LOCATION: (1)..(519)
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38 Met Val Asn Ser Asp Val Thr Arg His Ile Leu Ser Val Leu Val Gln
39 1               5                   10                  15
41 gac gta gac gat gac ttt tcc cgc gta tca ggt atg ttc acc cga cgc      96
42 Asp Val Asp Asp Asp Phe Ser Arg Val Ser Gly Met Phe Thr Arg Arg
43             20                  25                  30
45 gca ttc aac ctc gtg tcc ctc gtg tct gca aag acc gaa aca cac ggc     144
46 Ala Phe Asn Leu Val Ser Leu Val Ser Ala Lys Thr Glu Thr His Gly
47         35                  40                  45
49 atc aac cgc atc acg gtt gtt gtc gac gcc gac gag ctc aac att gag     192
50 Ile Asn Arg Ile Thr Val Val Val Asp Ala Asp Glu Leu Asn Ile Glu
51     50                  55                  60
53 cag atc aac aag cag ctc aac aag ctg atc ccc gtg ctc aaa gtc gtg     240
54 Gln Ile Asn Lys Gln Leu Asn Lys Leu Ile Pro Val Leu Lys Val Val
55 65                  70                  75                  80
57 cga ctt gat gaa gag acc act atc gcc cgc gca atc atg ctg gtt aag     288
58 Arg Leu Asp Glu Glu Thr Thr Ile Ala Arg Ala Ile Met Leu Val Lys
59             85                  90                  95
61 gtc tct gcg gac agc acc aac cgt ccg cag atc gtc gac gcc gcg aac     336
62 Val Ser Ala Asp Ser Thr Asn Arg Pro Gln Ile Val Asp Ala Ala Asn
63             100                 105                 110
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65 atc ttc cgc gcc cga gtc gtc gac gtg gct cca gac tct gtg gtt att       384
66 Ile Phe Arg Ala Arg Val Val Asp Val Ala Pro Asp Ser Val Val Ile
67     115             120             125
69 gaa tcc aca ggc acc cca ggc aag ctc cgc gca ctg ctt gac gtg atg       432
70 Glu Ser Thr Gly Thr Pro Gly Lys Leu Arg Ala Leu Leu Asp Val Met
71     130             135             140
73 gaa caa ttc gaa atc cgc gaa ctg atc caa tcc gga cag att gca ctc       480
74 Glu Gln Phe Glu Ile Arg Glu Leu Ile Gln Ser Gly Gln Ile Ala Leu
75 145             150             155             160
77 aac cgc ggt ccg aag acc atg gct ccg gcc aag atc taa                   519
78 Asn Arg Gly Pro Lys Thr Met Ala Pro Ala Lys Ile
79             165             170
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88 <223> OTHER INFORMATION: Synthetic Construct
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100 Ala Phe Asn Leu Val Ser Leu Val Ser Ala Lys Thr Glu Thr His Gly
101         35              40              45
104 Ile Asn Arg Ile Thr Val Val Val Asp Ala Asp Glu Leu Asn Ile Glu
105     50              55              60
108 Gln Ile Asn Lys Gln Leu Asn Lys Leu Ile Pro Val Leu Lys Val Val
109 65              70              75              80
112 Arg Leu Asp Glu Glu Thr Thr Ile Ala Arg Ala Ile Met Leu Val Lys
113             85              90              95
116 Val Ser Ala Asp Ser Thr Asn Arg Pro Gln Ile Val Asp Ala Ala Asn
117             100             105             110
120 Ile Phe Arg Ala Arg Val Val Asp Val Ala Pro Asp Ser Val Val Ile
121         115             120             125
124 Glu Ser Thr Gly Thr Pro Gly Lys Leu Arg Ala Leu Leu Asp Val Met
125     130             135             140
128 Glu Gln Phe Glu Ile Arg Glu Leu Ile Gln Ser Gly Gln Ile Ala Leu
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154 gac gta gac ggt gac ttt tcc cgc gta tca ggt atg ttc acc cga cgc     96
155 Asp Val Asp Gly Asp Phe Ser Arg Val Ser Gly Met Phe Thr Arg Arg
156             20                  25                  30
158 gca ttc aac ctc gtg tcc ctc gtg tct gca aag acc gaa aca cac ggc    144
159 Ala Phe Asn Leu Val Ser Leu Val Ser Ala Lys Thr Glu Thr His Gly
160         35                  40                  45
162 atc aac cgc atc acg gtt gtt gtc gac gcc gac gag ctc aac att gag    192
163 Ile Asn Arg Ile Thr Val Val Val Asp Ala Asp Glu Leu Asn Ile Glu
164     50                  55                  60
166 cag atc acc aag cag ctc aac aag ctg atc ccc gtg ctc aaa gtc gtg    240
167 Gln Ile Thr Lys Gln Leu Asn Lys Leu Ile Pro Val Leu Lys Val Val
168 65                  70                  75                  80
170 cga ctt gat gaa gag acc act atc gcc cgc gca atc atg ctg gtt aag    288
171 Arg Leu Asp Glu Glu Thr Thr Ile Ala Arg Ala Ile Met Leu Val Lys
172             85                  90                  95
174 gtc tct gcg gac agc acc aac cgt ccg cag atc gtc gac gcc gcg aac    336
175 Val Ser Ala Asp Ser Thr Asn Arg Pro Gln Ile Val Asp Ala Ala Asn
176             100                 105                 110
178 atc ttc cgc gcc cga gtc gtc gac gtg gct cca gac tct gtg gtt att    384
179 Ile Phe Arg Ala Arg Val Val Asp Val Ala Pro Asp Ser Val Val Ile
180         115                 120                 125
182 gaa tcc aca ggc acc cca ggc aag ctc cgc gca ctg ctt gac gtg atg    432
183 Glu Ser Thr Gly Thr Pro Gly Lys Leu Arg Ala Leu Leu Asp Val Met
184     130                 135                 140
186 gaa cca tcc gga atc gcg gaa ctg atc caa tcc gga cag att gca ctc    480
187 Glu Pro Ser Gly Ile Ala Glu Leu Ile Gln Ser Gly Gln Ile Ala Leu
188 145                 150                 155                 160
190 aac cgc ggt ccg aag acc atg gct ccg gcc aag atc taa               519
191 Asn Arg Gly Pro Lys Thr Met Ala Pro Ala Lys Ile
192                 165                 170
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196 <211> LENGTH: 172
197 <212> TYPE: PRT
198 <213> ORGANISM: Artificial Sequence
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201 <223> OTHER INFORMATION: Synthetic Construct
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210             20                  25                  30
213 Ala Phe Asn Leu Val Ser Leu Val Ser Ala Lys Thr Glu Thr His Gly
214         35                  40                  45
217 Ile Asn Arg Ile Thr Val Val Val Asp Ala Asp Glu Leu Asn Ile Glu
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218      50                      55                    60
221 Gln Ile Thr Lys Gln Leu Asn Lys Leu Ile Pro Val Leu Lys Val Val
222 65                      70            .       75                    80
225 Arg Leu Asp Glu Glu Thr Thr Ile Ala Arg Ala Ile Met Leu Val Lys
226                 85                      90                    95
229 Val Ser Ala Asp Ser Thr Asn Arg Pro Gln Ile Val Asp Ala Ala Asn
230                 100                   105                   110
233 Ile Phe Arg Ala Arg Val Val Asp Val Ala Pro Asp Ser Val Val Ile
234         115                   120           .       125
237 Glu Ser Thr Gly Thr Pro Gly Lys Leu Arg Ala Leu Leu Asp Val Met
238     130                   135                   140
241 Glu Pro Ser Gly Ile Ala Glu Leu Ile Gln Ser Gly Gln Ile Ala Leu
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246                 165                   170
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255 <223> OTHER INFORMATION: Synthetic DNA
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303 <223> OTHER INFORMATION: Synthetic DNA
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321             20                  25                  30
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329     50                  55
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334 <212> TYPE: PRT
335 <213> ORGANISM: S. cinnamonensis
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367             20                  25                  30
370 Ala Phe Asn Val Glu Gly Ile Leu Cys Leu Pro Ile Gln Asp Ser Asp
371         35                  40                  45
374 Lys Ser His Ile Trp Leu Leu Val Asn
375     50                  55
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VERIFICATION SUMMARY   DATE: 03/28/2007
PATENT APPLICATION: **US/10/561,906**   TIME: 14:53:00

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